

STIC Search Report Biotech-Chem Library

STIC Database Tracking Number: 1977/94

TO: Christian Fronda Location: rem-2d78/2c70

Art Unit: 1652

Wednesday, August 09, 2006

Case Serial Number: 09/435770

From: Kristine Hensle

Location: Biotech-Chem Library

REM-1A70

Phone: (571)272-4161

Kristine.Hensle@uspto.gov

Search Notes

Examiner Fronda,

See attached results.

If you have any questions about this search feel free to contact me at any time.

Thank you for using STIC search services!

Kristine Hensle Reference Librarian STIC Biotech/Chem Library (571)272-4161



STIC-Biotech/ChemLib

197774

From:

Chan, Christina

Sent:

Tuesday, August 08, 2006 8:19 AM Fronda, Christian; STIC-Biotech/ChemLib

To: Subject:

RE: Rush Search for Serial No. 09/435,770

Please rush. Thanks Chris

Chris Chan

TC 1600 New Hire Training Coordinator and SPE 1644 (571)-272-0841 Remsen, 3E89

----Original Message-----

From:

Fronda, Christian

Sent:

Monday, August 07, 2006 6:38 PM

To:

Chan, Christina

Subject:

Rush Search for Serial No. 09/435,770

Importance: High

I would like to request a Rush Search for Serial No. 09/435,770 as listed below since it is an amended case. Thank you.

Christian L. Fronda Art Unit 1652 Office REM 2D78 Mailbox REM 2C70 (571)272-0929

Please perform sequence and interference search for Serial No. 09/435,770

1. Please search SEQ ID Nos: 1-6 against amino acid commercial, PGPub, issued databases, and interference databases.

Please save on **COMPUTER DISKETTES**.

Please save the interference searches on a separate disk.

Thank you very much.

Christian L. Fronda Art Unit 1652 Office REM 2D78 Mailbox REM 2C70 (571)272-0929

Searcher:	
Searcher Phone:	
Date Searcher Picked up:	
Date completed:	
Searcher Prep Time:	
Online Time:	

Type	of Search
NA#	_ AA#:
S/L: C	ligomer:
Encode/Tran	sl:
Structure #:_	Text:
Inventor:	Litigation:

Vendors and cost where applicable
STN:
DIALOG:
QUESTEL/ORBIT:
LEXIS/NEXIS:
SEQUENCE SYSTEM:
WWW/Internet:
Other (Specify):



STIC SEARCH RESULTS FEEDBACK FORM

Biotech-Chem Library

Questions about the scope or the results of the search? Contact the searcher or contact:

Mary Hale, Information Branch Supervisor 571-272-2507 Remsen E01 D86

VOI	untary Results Feedback Form
>	I am an examiner in Workgroup: Example: 1610
\triangleright	Relevant prior art found, search results used as follows:
	☐ 102 rejection
	☐ 103 rejection
	Cited as being of interest.
	Helped examiner better understand the invention.
	Helped examiner better understand the state of the art in their technology.
	Types of relevant prior art found:
	☐ Foreign Patent(s)
ε	 Non-Patent Literature (journal articles, conference proceedings, new product announcements etc.)
>	Relevant prior art not found:
•	Results verified the lack of relevant prior art (helped determine patentability).
-	Results were not useful in determining patentability or understanding the invention.
Со	mments:

Drop off or send completed forms to STIC/Blotech-Chem Library Remson Eldg.



Protein Sequence Searches - February 2005

All of the sequence databases on ABSS have recently been updated.

- Please note that the curators of the UniProt database have purged some temporary accession numbers from the most recent version of UniProt. These sequences have been assigned new permanent accession numbers. The new UniProt record may not contain the previous temporary accession number.
- If you encounter an accession number from an older search run against UniProt (results file extension .rup) that can no longer be found in the database, the permanent record with the new accession number can be found by searching the old accession number in the UniProt Protein Archive database (UniPARC) at:

http://www.pir.uniprot.org/database/archive.shtml

If you have any questions regarding this information or your results, please contact any STIC searcher.

When submitting sequence search results for scanning into IFW, please include a copy of this attachment to assist any future Examiners or members of the public who may encounter UniProt temporary accession numbers.

November 2005

Published_Applications Nucleic Acid and Published_Applications Amino Acid database searches now generate two sets of results each. The Published_Applications databases have been split into two parts to reduce the amount of time required for their daily updates. This results in more machine time being available for processing searches.

Newly published applications will appear in the Published_Applications_New databases: older published applications make up the Published_Applications_Main databases.

Searches run against Nucleic Acid Published_Applications produce two sets of results, with the extensions .rnpbm (Published_Applications_NA_Main) and .rnpbm (Published_Applications_NA_New).

Searches run against Amino Acid Published_Applications produce two sets of results, with the extensions .rapbm (Published_Applications_AA_New).

Pending Nucleic Acid and Pending Amino Acid database searches generate two sets of results each. The Pending databases have been split into two parts to reduce the amount of time required for their daily updates. This results in more machine time being available for processing searches.

Searches run against the Nucleic Acid Pending database produce two sets of results, with the extensions .rnpm and .rnpn

Searches run against the Amino Acid Pending database produce two sets of results, with the extensions .rapm and .rapn

Because they contain data that is confidential, the results of Pending database searches should not be left in the case.